

**10/527725**

**RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.

Application Serial Number: 10/527,725

Source: PCT

Date Processed by STIC: 3-25-05

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PCT

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/10/527,725**

**DATE: 03/25/2005**  
**TIME: 08:58:09**

**Input Set : E:\SEQID.ST25.txt**  
**Output Set: N:\CRF4\03252005\J527725.raw**

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3 <110> APPLICANT: Texas Tech University
4     Shaw, Robert W.
6 <120> TITLE OF INVENTION: Inhibition of metallo-beta-lactamase
8 <130> FILE REFERENCE: 218601.00007
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/527,725
C--> 10 <141> CURRENT FILING DATE: 2005-03-14
10 <160> NUMBER OF SEQ ID NOS: 6
12 <170> SOFTWARE: PatentIn version 3.3
14 <210> SEQ ID NO: 1
15 <211> LENGTH: 16
16 <212> TYPE: DNA
17 <213> ORGANISM: artificial sequence
19 <220> FEATURE:
20 <223> OTHER INFORMATION: A 16 mer with a NdeI restriction site for SELEX.
22 <400> SEQUENCE: 1
23 ggcggccatatg cgccgcg                                     16
26 <210> SEQ ID NO: 2
27 <211> LENGTH: 15
28 <212> TYPE: DNA
29 <213> ORGANISM: artificial sequence
31 <220> FEATURE:
32 <223> OTHER INFORMATION: A 15 mer with a SacI restriction site for SELEX.
34 <400> SEQUENCE: 2
35 cgcgagctcc ggcgcg                                      15
38 <210> SEQ ID NO: 3
39 <211> LENGTH: 30
40 <212> TYPE: DNA
41 <213> ORGANISM: artificial sequence
43 <220> FEATURE:
44 <223> OTHER INFORMATION: This is the aptamer sequence after 16 rounds of SELEX.
47 <220> FEATURE:
48 <221> NAME/KEY: misc_feature
49 <222> LOCATION: (2)..(2)
50 <223> OTHER INFORMATION: n is A or T or G or C
52 <220> FEATURE:
53 <221> NAME/KEY: misc_feature
54 <222> LOCATION: (4)..(4)
55 <223> OTHER INFORMATION: n is A or T or G or C
57 <220> FEATURE:
58 <221> NAME/KEY: misc_feature
59 <222> LOCATION: (6)..(6)
60 <223> OTHER INFORMATION: N is A or T or G or C
62 <220> FEATURE:

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63 <221> NAME/KEY: misc\_feature  
64 <222> LOCATION: (7)..(7)  
65 <223> OTHER INFORMATION: N is A or T or G or C  
67 <220> FEATURE:  
68 <221> NAME/KEY: misc\_feature  
69 <222> LOCATION: (8)..(8)  
70 <223> OTHER INFORMATION: N is A or T or G or C  
72 <220> FEATURE:  
73 <221> NAME/KEY: misc\_feature  
74 <222> LOCATION: (11)..(11)  
75 <223> OTHER INFORMATION: N is A or T or G or C  
77 <220> FEATURE:  
78 <221> NAME/KEY: misc\_feature  
79 <222> LOCATION: (12)..(12)  
80 <223> OTHER INFORMATION: N is A or T or G or C  
82 <220> FEATURE:  
83 <221> NAME/KEY: misc\_feature  
84 <222> LOCATION: (13)..(13)  
85 <223> OTHER INFORMATION: N is A or T or G or C  
87 <220> FEATURE:  
88 <221> NAME/KEY: misc\_feature  
89 <222> LOCATION: (15)..(15)  
90 <223> OTHER INFORMATION: N is A or T or G or C  
92 <220> FEATURE:  
93 <221> NAME/KEY: misc\_feature  
94 <222> LOCATION: (17)..(17)  
95 <223> OTHER INFORMATION: N is A or T or G or C  
97 <220> FEATURE:  
98 <221> NAME/KEY: misc\_feature  
99 <222> LOCATION: (18)..(18)  
100 <223> OTHER INFORMATION: N is A or T or G or C  
102 <220> FEATURE:  
103 <221> NAME/KEY: misc\_feature  
104 <222> LOCATION: (20)..(20)  
105 <223> OTHER INFORMATION: N is A or T or G or C  
107 <220> FEATURE:  
108 <221> NAME/KEY: misc\_feature  
109 <222> LOCATION: (21)..(21)  
110 <223> OTHER INFORMATION: N is A or T or G or C  
112 <220> FEATURE:  
113 <221> NAME/KEY: misc\_feature  
114 <222> LOCATION: (25)..(25)  
115 <223> OTHER INFORMATION: N is A or T or G or C  
117 <220> FEATURE:  
118 <221> NAME/KEY: misc\_feature  
119 <222> LOCATION: (26)..(26)  
120 <223> OTHER INFORMATION: N is A or T or G or C  
122 <220> FEATURE:  
123 <221> NAME/KEY: misc\_feature

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124 <222> LOCATION: (27)..(27)  
 125 <223> OTHER INFORMATION: N is A or T or G or C  
 127 <220> FEATURE:  
 128 <221> NAME/KEY: misc\_feature  
 129 <222> LOCATION: (28)..(28)  
 130 <223> OTHER INFORMATION: N is A or T or G or C  
 132 <400> SEQUENCE: 3  
**W--> 133 ancnannnntt nnntngnngn ncatnnnnaa** 30  
 136 <210> SEQ ID NO: 4  
 137 <211> LENGTH: 30  
 138 <212> TYPE: DNA  
 139 <213> ORGANISM: artificial sequence  
 141 <220> FEATURE:  
 142 <223> OTHER INFORMATION: This is the aptamer 30-mer sequence after 21 rounds of SELEX.  
 144 <400> SEQUENCE: 4  
 145 aacccaaactt ggatcggtgc acatgtcgaa 30  
 148 <210> SEQ ID NO: 5  
 149 <211> LENGTH: 10  
 150 <212> TYPE: DNA  
 151 <213> ORGANISM: artificial sequence  
 153 <220> FEATURE:  
 154 <223> OTHER INFORMATION: This is a 10 mer that contains a specific stem loop structure 10  
 157 <220> FEATURE:  
 158 <221> NAME/KEY: stem\_loop  
 159 <222> LOCATION: (1)..(10)  
 161 <400> SEQUENCE: 5  
 162 cccaaacttgg 10  
 165 <210> SEQ ID NO: 6  
 166 <211> LENGTH: 61  
 167 <212> TYPE: DNA  
 168 <213> ORGANISM: artificial sequence  
 170 <220> FEATURE:  
 171 <223> OTHER INFORMATION: the aptamer (61-mer)  
 173 <400> SEQUENCE: 6  
 174 gcgccatatg cgcgcaacc aaacttggat cggcgcacat gtcgaacgcg cggagctcgc 60  
 176 g 61

RAW SEQUENCE LISTING ERROR SUMMARY                   DATE: 03/25/2005  
PATENT APPLICATION: US/10/527,725                   TIME: 08:58:10

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; N Pos. 2, 4, 6, 7, 8, 11, 12, 13, 15, 17, 18, 20, 21, 25, 26, 27, 28

**VERIFICATION SUMMARY**  
PATENT APPLICATION: US/10/527,725

DATE: 03/25/2005  
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Input Set : E:\SEQID.ST25.txt  
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L:10 M:270 C: Current Application Number differs, Replaced Current Application No  
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:133 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0